

Prelim. Amdt dated November 21, 2003
ANNOTATED SHEETS SHOWING CHANGES-SEQUENCE LISTING

therapeutically active agent may be associated with a container holding a flowable carrier component (e.g. a container may hold fibrinogen and C3).

5 SEQUENCE LISTING

(i) GENERAL INFORMATION:

10 (i) APPLICANT: LISA MCKERRACHER

(ii) TITLE OF INVENTION: Methods for making and delivering Rho-antagonist
tissue adhesive formulations to the injured mammalian
central and peripheral nervous systems and uses thereof

15 (iii) NUMBER OF SEQUENCES: 3

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: BROULLETTE KOSIE

20 (B) STREET: 1100 RENE LESVEQUE BLVD WEST

(C) PROV/STATE: QUEBEC

(D) COUNTRY: CANADA

(E) POSTAL/ZIP CODE: H3B 5C9

25 (v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: Floppy disk

(B) COMPUTER: IBM PC compatible

(C) OPERATING SYSTEM: PC DOS/MS-DOS

30 (D) SOFTWARE: ASCII (TEXT) Patent In version 3.1

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER:

~~(B) FILING DATE:~~

~~(C) CLASSIFICATION:~~

~~(vii) ATTORNEY/AGENT INFORMATION:~~

5 ~~(A) NAME: RONALD S. KOSIE~~

~~(B) REGISTRATION NO.: 28,814~~

~~(C) REFERENCE/DOCKET NO.: 06447-003-US-2~~ 06447-011

~~(D) TEL. NO.: (514) 397-8500~~

~~(E) FAX NO.: (514) 397-8515~~

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(2) INFORMATION FOR SEQ ID NO: 1:

(i) SEQUENCE CHARACTERISTICS:

15 (A) LENGTH:

(B) TYPE:

(C) STRANDEDNESS:

(D) TOPOLOGY:

20 (iii) MOLECULE TYPE:

(v) FRAGMENT TYPE:

25 ~~(vi) ORIGINAL SOURCE:~~

(A) ORGANISM:

~~(vii) IMMEDIATE SOURCE:~~

30 ~~(ix) FEATURE:~~

(A) NAME/KEY:

~~(B) LOCATION:~~

~~(D) OTHER INFORMATION:~~

5 (x) PUBLICATION INFORMATION:

(A) AUTHORS:

(B) TITLE:

(C) JOURNAL:

~~(D) VOLUME:~~

10 (E) ISSUE:

(F) PAGES:

(G) DATE:

(H) DOCUMENT NO.:

(I) FILING DATE:

15 (J) PUBLICATION DATE:

(K) RELEVANT RESIDUES IN SEQ ID NO:

(x) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

20 GTG GCG ACC CTT CCC AAA TCG GAT CTG GTT CCG CGT GGA TCC TCT AGA
 5 10 15
GTC GAC CTG CAG GCA TGC AAT GCT TAT TCC ATT AAT CAA AAG GCT TAT
 20 25 30
TCA AAT ACT TAC CAG GAG TTT ACT AAT ATT GAT CAA GCA AAA GCT TGG
25 35 40 45
GGT AAT GCT CAG TAT AAA AAG TAT GGA CTA AGC AAA TCA GAA AAA GAA
 50 55 60
GCT ATA GTA TCA TAT ACT AAA AGC GCT AGT GAA ATA AAT GGA AAG CTA
 65 70 75 80
30 AGA CAA AAT AAG GGA GTT ATC AAT GGA TTT CCT TCA AAT TTA ATA AAA
 85 90 95
CAA GTT GAA CTT TTA GAT AAA TCT TTT AAT AAA ATG AAG ACC CCT GAA

~~100~~ ~~105~~ ~~110~~

AAT ATT ATG TTA TTT AGA GGC GAC GAC CCT GCT TAT TTA GGA ACA GAA

~~115~~ ~~120~~ ~~125~~

5 TTT CAA AAC ACT CTT CTT AAT TCA AAT GGT ACA ATT AAT AAA ACG GCT

~~130~~ ~~135~~ ~~140~~

TTT GAA AAG GCT AAA GCT AAG TTT TTA AAT AAA GAT AGA CTT GAA TAT

~~145~~ ~~150~~ ~~155~~ ~~160~~

GGA TAT ATT AGT ACT TCA TTA ATG AAT GTT TCT CAA TTT GCA GGA AGA

~~165~~ ~~170~~ ~~175~~

10 CCA ATT ATT ACA AAA TTT AAA GTA GCA AAA GGC TCA AAG GCA GGA TAT

~~180~~ ~~185~~ ~~190~~

ATT GAC CCT ATT AGT GCT TTT CAG GGA CAA CTT GAA ATG TTG CTT CCT

~~195~~ ~~200~~ ~~205~~

15 AGA CAT AGT ACT TAT CAT ATA GAC GAT ATG AGA TTG TCT TCT GAT GGT

~~210~~ ~~215~~ ~~220~~

AAA CAA ATA ATA ATT ACA GCA ACA ATG ATG GGC ACA GCT ATC AAT CCT

~~225~~ ~~230~~ ~~235~~ ~~240~~

AAA TAA

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25 (2) INFORMATION FOR SEQ ID NO: 2:

~~(i) SEQUENCE CHARACTERISTICS:~~

(A) LENGTH:

(B) TYPE:

~~(C) STRANDEDNESS:~~

~~(D) TOPOLOGY:~~

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~~(vi) ORIGINAL SOURCE:~~

(A) ORGANISM:

~~(ix) FEATURE:~~

~~(D) OTHER INFORMATION:~~

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~~(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:~~

10 GGATCCTCTA GAGTCGACCT GCAGGCATGC AATGCTTATT CCATTAATCA 50-
AAAGGCTTAT TCAAATACTT ACCAGGAGTT TACTAATATT GATCAAGCAA 100-
AAGCTTGGGG TAATGCTCAG TATAAAAAGT ATGGACTAAG CAAATCAGAA 150-
AAAGAAGCTA TAGTATCATA TACTAAAAGC GCTAGTGAAA TAAATGGAAA 200-
GCTAAGACAA AATAAGGGAG TTATCAATGG ATTCCTTCA AATTTAATAA 250-
15 AACAAGTTGA ACTTTTAGAT AAATCTTTTA ATAAAATGAA GACCCCTGAA 300-
AATATTATGT TATTAGAGG CGACGACCCT GCTTATTTAG GAACAGAATT 350-
TCAAACACT CTTCTTAATT CAAATGGTAC AATTAATAAA ACGGCTTTTG 400-
AAAAGGCTAA AGCTAAGTTT TTAAATAAAG ATAGACTTGA ATATGGATAT 450-
ATTAGTACTT CATTAATGAA TGTTTCTCAA TTTGCAGGAA GACCAATTAT 500-
20 TACAAAATTT AAAGTAGCAA AAGGCTCAA GGCAGGATAT ATTGACCCTA 550-
TTAGTGCTTT TCAGGGACAA CTTGAAATGT TGCTTCCTAG ACATAGTACT 600-
TATCATATAG ACGATATGAG ATTGTCTTCT GATGGTAAAC AAATAATAAT 650-
TACAGCAACA ATGATGGGCA CAGCTATCAA TCCTAAATAA

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(2) INFORMATION FOR SEQ ID NO: 3:

30 ~~(i) SEQUENCE CHARACTERISTICS:~~

(A) LENGTH:

(B) TYPE:

BEST AVAILABLE COPY

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(C) STRANDEDNESS:

(D) TOPOLOGY:

(vi) ORIGINAL SOURCE:

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(A) ORGANISM:

(ix) FEATURE:—

(D) OTHER INFORMATION:

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(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:

15

GSSRVDLQAC NAYSINQKAY SNTYQEFTNI DQAKAWGNAQ YKKYGLSKSE 50
KEAIVSYTKS ASEINGKLRQ NKGVINGFPS NLIKQVELLD KSFNKMKTPE 100
NIMLFXGDDP AYLGTEFQNT LLNSNGTINK TAFEKAKAKF LNXDRLEYGY 150
ISTSLMNVSQ FAGRPIITKF KVAKGSKAGY IDPISAFQGQ LEMLLPRHST 200
YHIDDMRLSS DGKQIIITAT MMGTAINPK

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